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1644

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TECH CENTER 1600/2900



1600

RAW SEQUENCE LISTING

DATE: 07/22/2002

PATENT APPLICATION: US/09/397,342A

TIME: 13:58:26

p#18

Input Set : A:\P1626R1.txt

Output Set: N:\CRF3\07222002\I397342A.raw

ENTERED

3 <110> APPLICANT: Adams, Sean
4 Pan, James
5 Zhong, Alan
7 <120> TITLE OF INVENTION: UCP4
9 <130> FILE REFERENCE: P1626R1
11 <140> CURRENT APPLICATION NUMBER: US 09/397,342A
12 <141> CURRENT FILING DATE: 1999-09-15
14 <150> PRIOR APPLICATION NUMBER: US 60/101,279
15 <151> PRIOR FILING DATE: 1998-09-22
17 <150> PRIOR APPLICATION NUMBER: US 60/114,223
18 <151> PRIOR FILING DATE: 1998-12-30
20 <150> PRIOR APPLICATION NUMBER: US 60/129,674
21 <151> PRIOR FILING DATE: 1999-04-16
23 <160> NUMBER OF SEQ ID NOS: 18
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 323
27 <212> TYPE: PRT
28 <213> ORGANISM: Homo sapiens
30 <400> SEQUENCE: 1
31 Met Ser Val Pro Glu Glu Glu Arg Leu Leu Pro Leu Thr Gln
32 1 5 10 15
34 Arg Trp Pro Arg Ala Ser Lys Phe Leu Leu Ser Gly Cys Ala Ala
35 20 25 30
37 Thr Val Ala Glu Leu Ala Thr Phe Pro Leu Asp Leu Thr Lys Thr
38 35 40 45
40 Arg Leu Gln Met Gln Gly Glu Ala Ala Leu Ala Arg Leu Gly Asp
41 50 55 60
43 Gly Ala Arg Glu Ser Ala Pro Tyr Arg Gly Met Val Arg Thr Ala
44 65 70 75
46 Leu Gly Ile Ile Glu Glu Glu Gly Phe Leu Lys Leu Trp Gln Gly
47 80 85 90
49 Val Thr Pro Ala Ile Tyr Arg His Val Val Tyr Ser Gly Gly Arg
50 95 100 105
52 Met Val Thr Tyr Glu His Leu Arg Glu Val Val Phe Gly Lys Ser
53 110 115 120
55 Glu Asp Glu His Tyr Pro Leu Trp Lys Ser Val Ile Gly Gly Met
56 125 130 135
58 Met Ala Gly Val Ile Gly Gln Phe Leu Ala Asn Pro Thr Asp Leu
59 140 145 150
61 Val Lys Val Gln Met Gln Met Glu Gly Lys Arg Lys Leu Glu Gly
62 155 160 165
64 Lys Pro Leu Arg Phe Arg Gly Val His His Ala Phe Ala Lys Ile
65 170 175 180

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```

67  Leu Ala Glu Gly Gly Ile Arg Gly Leu Trp Ala Gly Trp Val Pro
68                      185                      190                      195
70  Asn Ile Gln Arg Ala Ala Leu Val Asn Met Gly Asp Leu Thr Thr
71                      200                      205                      210
73  Tyr Asp Thr Val Lys His Tyr Leu Val Leu Asn Thr Pro Leu Glu
74                      215                      220                      225
76  Asp Asn Ile Met Thr His Gly Leu Ser Ser Leu Cys Ser Gly Leu
77                      230                      235                      240
79  Val Ala Ser Ile Leu Gly Thr Pro Ala Asp Val Ile Lys Ser Arg
80                      245                      250                      255
82  Ile Met Asn Gln Pro Arg Asp Lys Gln Gly Arg Gly Leu Leu Tyr
83                      260                      265                      270
85  Lys Ser Ser Thr Asp Cys Leu Ile Gln Ala Val Gln Gly Glu Gly
86                      275                      280                      285
88  Phe Met Ser Leu Tyr Lys Gly Phe Leu Pro Ser Trp Leu Arg Met
89                      290                      295                      300
91  Thr Pro Trp Ser Met Val Phe Trp Leu Thr Tyr Glu Lys Ile Arg
92                      305                      310                      315
94  Glu Met Ser Gly Val Ser Pro Phe
95                      320

```

97 <210> SEQ ID NO: 2

98 <211> LENGTH: 1039

99 <212> TYPE: DNA

100 <213> ORGANISM: Homo sapiens

102 <400> SEQUENCE: 2

```

103  ccgagctcgg atcccgttat cgtcttgccg tactgctgaa tgtccgtccc 50
105  ggaggaggag gagaggcttt tgccgctgac ccagagatgg ccccgagcga 100
107  gcaaattcct actgtccggc tgcgcggcta ccgtggccga gctagcaacc 150
109  tttcccctgg atctcacaaa aactcgactc caaatgcaag gagaagcagc 200
111  tcttgctcgg ttgggagacg gtgcaagaga atctgcccc tataggggaa 250
113  tgggtgcgcac agccctaggg atcattgaag aggaaggctt tctaaagctt 300
115  tggcaaggag tgacaccgcg catttacaga cacgtagtgt attctggagg 350
117  tcgaatggtc acatatgaac atctccgaga ggttggtgtt ggcaaaagtg 400
119  aagatgagca ttatcccctt tggaaatcag tcattggagg gatgatggct 450
121  ggtgtttatt gccagttttt agccaatcca actgacctag tgaaggttca 500
123  gatgcaaatt gaaggaaaaa ggaaactgga aggaaaacca ttgcgatttc 550
125  gtggtgtaca tcatgcattt gcaaaaatct tagctgaagg aggaatacga 600
127  gggctttggg caggctgggt acccaatata caaagagcag cactggtgaa 650
129  tatgggagat ttaaccactt atgatacagt gaaacactac ttggtattga 700
131  atacaccact tgaggacaat atcatgactc acggtttatc aagtttatgt 750
133  tctggactgg tagcttctat tctgggaaca ccagccgatg tcatcaaaag 800
135  cagaataatg aatcaaccac gagataaaca aggaagggga cttttgtata 850
137  aatcatcgac tgactgcttg attcaggctg ttcaagggtg aggattcatg 900
139  agtctatata aaggcttttt accatcttgg ctgagaatga ccccttggtc 950
141  aatggtgttc tggttactt atgaaaaaat cagagagatg agtggagtca 1000
143  gtccatttta agaattctgc agatatccat cacactggc 1039

```

145 <210> SEQ ID NO: 3

146 <211> LENGTH: 31

147 <212> TYPE: DNA

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```

C--> 148 <213> ORGANISM: Artificial
      150 <220> FEATURE:
W--> 151 <221> NAME/KEY: Misc-feature
      152 <222> LOCATION: 1-31
      153 <223> OTHER INFORMATION: Sequence is synthesized
      155 <400> SEQUENCE: 3
      156   cgcggaatccc gttatcgtct tgcgctactg c 31
      158 <210> SEQ ID NO: 4
      159 <211> LENGTH: 34
      160 <212> TYPE: DNA
C--> 161 <213> ORGANISM: Artificial
      163 <220> FEATURE:
W--> 164 <221> NAME/KEY: Misc-feature
      165 <222> LOCATION: 1-34
      166 <223> OTHER INFORMATION: Sequence is synthesized
      168 <400> SEQUENCE: 4
      169   gcggaattct taaaatggac tgactccact catc 34
      171 <210> SEQ ID NO: 5
      172 <211> LENGTH: 1248
      173 <212> TYPE: DNA
C--> 174 <213> ORGANISM: Artificial
      176 <220> FEATURE:
W--> 177 <221> NAME/KEY: Misc-feature
      178 <222> LOCATION: 1-1248
      179 <223> OTHER INFORMATION: Sequence is synthesized
      181 <220> FEATURE:
      182 <221> NAME/KEY: unsure
      183 <222> LOCATION: 1231
      184 <223> OTHER INFORMATION: unknown base
      186 <400> SEQUENCE: 5
      187   cgttatcgtc ttgcgctact gctgaatgtc cgtcccggag gaggaggaga 50
      189   ggcttttgcc gctgaccag agatggcccc gagcgagcaa attcctactg 100
      191   tccggtgcg cggtaccgt ggccgagcta gcaacctttc ccttggtatc 150
      193   cacaaaaact cgactccaaa tgcaaggaga agcagctctt gctcggttgg 200
      195   gagacggtgc aagagaatct gccccctata ggggaatggg gcgcacagcc 250
      197   ctagggatca ttgaagagga aggttttcta aagctttggc aaggagtgc 300
      199   acccgccatt tacagacacg tagttatttc tggaggtcga atggtcacat 350
      201   atgaacatct ccgagaggtt gtgtttggca aaagtgaaga tgagcattat 400
      203   cccctttgga aatcagtcac tggagggatg atggctggtg ttattggcca 450
      205   gtttttagcc aatccaactg acctagttaa gggttcagatg caaatggaag 500
      207   gaaaaaggaa actggaagga aaaccattgc gatttcgtgg tgtacatcat 550
      209   gcatttgcaa aaatcttagc tgaaggagga atacgaaggc tttgggcagg 600
      211   ctgggtaccc aatatacaaa gagcagcact ggtgaatatg ggagatttaa 650
      213   ccacttatga tacagtgaac cactacttgg tattgaatac accacttgag 700
      215   gacaatatca tgactcacgg ttatcaagt ttatgttctg gactggtagc 750
      217   ttctattctg ggaacaccag ccgatgtcat caaaagcaga ataataatc 800
      219   aaccacgaga taaacaagga aggggacttt tgtataaatc atcgactgac 850
      221   tgcttgattc aggtgttca aggtgaagga ttcattgagtc tatataaagg 900
      223   ctttttacca tcttggctga gaatgacccc ttggtcaatg gtgttctggc 950

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Input Set : A:\P1626R1.txt

Output Set: N:\CRF3\07222002\I397342A.raw

```

225 ttacttatga aaaaatcaga gagatgagtg gagtcagtcc attttaaac 1000
227 cctaaagatg caacccttaa agatacagtg ttcagtatta ttgaaatatg 1050
229 ggcattctgca acacataccc cctattatct ctacctcttt aggaagacac 1100
231 ctattccaca gagactgatt tatagggggc agcactttat ttttttctgg 1150
233 aaacccaagt tctctttgac tcctcttttt gtccaaaagt gatctggtcg 1200
W--> 235 gatctcaciaa ggccatccaa tgagaccccg nacagcattt tctaaaga 1248
237 <210> SEQ ID NO: 6
238 <211> LENGTH: 58
239 <212> TYPE: DNA
C--> 240 <213> ORGANISM: Artificial
242 <220> FEATURE:
W--> 243 <221> NAME/KEY: Misc-feature
244 <222> LOCATION: 1-58
245 <223> OTHER INFORMATION: Sequence is synthesized
247 <400> SEQUENCE: 6
248 cgcggatccg aaatggacta caaggacgac gatgacaagt ccgtcccga 50
250 ggaggagg 58
252 <210> SEQ ID NO: 7
253 <211> LENGTH: 35
254 <212> TYPE: DNA
C--> 255 <213> ORGANISM: Artificial
257 <220> FEATURE:
W--> 258 <221> NAME/KEY: Misc-feature
259 <222> LOCATION: 1-35
260 <223> OTHER INFORMATION: Sequence is synthesized
262 <400> SEQUENCE: 7
263 gcgaagcttg ccatgggttg actgaagcct tcaga 35
265 <210> SEQ ID NO: 8
266 <211> LENGTH: 33
267 <212> TYPE: DNA
C--> 268 <213> ORGANISM: Artificial
270 <220> FEATURE:
W--> 271 <221> NAME/KEY: Misc-feature
272 <222> LOCATION: 1-33
273 <223> OTHER INFORMATION: Sequence is synthesized
275 <400> SEQUENCE: 8
276 cgcgaattct caaaacggtg attcccgtaa cat 33
278 <210> SEQ ID NO: 9
279 <211> LENGTH: 61
280 <212> TYPE: DNA
C--> 281 <213> ORGANISM: Artificial
283 <220> FEATURE:
W--> 284 <221> NAME/KEY: Misc-feature
285 <222> LOCATION: 1-61
286 <223> OTHER INFORMATION: Sequence is synthesized
288 <400> SEQUENCE: 9
289 gcgaagcttg ccatggacta caaggacgac gatgacaagg ttggactgaa 50
291 gccttcagac g 61
293 <210> SEQ ID NO: 10

```

RAW SEQUENCE LISTING

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Input Set : A:\P1626R1.txt

Output Set: N:\CRF3\07222002\I397342A.raw

```

294 <211> LENGTH: 19
295 <212> TYPE: DNA
C--> 296 <213> ORGANISM: Artificial
298 <220> FEATURE:
W--> 299 <221> NAME/KEY: Misc-feature
300 <222> LOCATION: 1-19
301 <223> OTHER INFORMATION: Sequence is synthesized
303 <400> SEQUENCE: 10
304 aatgcctatc gccgaggag 19
306 <210> SEQ ID NO: 11
307 <211> LENGTH: 20
308 <212> TYPE: DNA
C--> 309 <213> ORGANISM: Artificial
311 <220> FEATURE:
W--> 312 <221> NAME/KEY: Misc-feature
313 <222> LOCATION: 1-20
314 <223> OTHER INFORMATION: Sequence is synthesized
316 <400> SEQUENCE: 11
317 gtaggaactt gctcgtccgg 20
319 <210> SEQ ID NO: 12
320 <211> LENGTH: 22
321 <212> TYPE: DNA
C--> 322 <213> ORGANISM: Artificial
324 <220> FEATURE:
W--> 325 <221> NAME/KEY: Misc-feature
326 <222> LOCATION: 1-22
327 <223> OTHER INFORMATION: Sequence is synthesized
329 <400> SEQUENCE: 12
330 tgctcgcgct caccgagaga tg 22
332 <210> SEQ ID NO: 13
333 <211> LENGTH: 24
334 <212> TYPE: DNA
C--> 335 <213> ORGANISM: Artificial
337 <220> FEATURE:
W--> 338 <221> NAME/KEY: Misc-feature
339 <222> LOCATION: 1-24
340 <223> OTHER INFORMATION: Sequence is synthesized
342 <400> SEQUENCE: 13
343 gaaatcgtgc gtgacatcaa agag 24
345 <210> SEQ ID NO: 14
346 <211> LENGTH: 23
347 <212> TYPE: DNA
C--> 348 <213> ORGANISM: Artificial
350 <220> FEATURE:
W--> 351 <221> NAME/KEY: Misc-feature
352 <222> LOCATION: 1-23
353 <223> OTHER INFORMATION: Sequence is synthesized
355 <400> SEQUENCE: 14
356 ctcttctgc atcctgtcag caa 23

```

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/397,342A

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 1231

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:3,4,5,6,7,8,9,10,11,12,13,14,15